



0590  
05-28

## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

#13

Application Serial Number: 09/718,321  
Source: OIP  
Date Processed by STIC: 5/21/2002

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

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Revised 01/29/2002

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OIPE

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/718,321

DATE: 05/21/2002  
 TIME: 14:28:49

Input Set : A:\Cura-99.app  
 Output Set: N:\CRF3\05212002\I718321.raw

**Does Not Comply  
 Corrected Diskette Needed**

4 <110> APPLICANT: Shimkets, Richard A.  
 5 Leach, Martin D.  
 7 <120> TITLE OF INVENTION: NUCLEIC ACIDS CONTAINING SINGLE NUCLEIC ACID POLYMORPHISMS  
 AND METHODS OF  
 8 USE THEREOF  
 10 <130> FILE REFERENCE: 15966-599  
 12 <140> CURRENT APPLICATION NUMBER: US/09/718,321  
 12 <141> CURRENT FILING DATE: 2002-05-07  
 12 <150> PRIOR APPLICATION NUMBER: 60/163,783  
 13 <151> PRIOR FILING DATE: 1999-11-24  
 15 <160> NUMBER OF SEQ ID NOS: 1468  
 17 <170> SOFTWARE: CuraGen Patent Formatter Version 0.9

## ERRORED SEQUENCES

23675 <210> SEQ ID NO: 1405  
 23676 <211> LENGTH: 12 *11 shown*  
 23677 <212> TYPE: PRT  
 23678 <213> ORGANISM: Homo sapiens  
 23680 <220> FEATURE:  
 23681 <221> NAME/KEY: VARIANT  
 23682 <222> LOCATION: (7)...(0)  
 23683 <223> OTHER INFORMATION: cSNP translation  
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 23767 <223> OTHER INFORMATION: cSNP translation  
 23769 <400> SEQUENCE: 1411  
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 E--> 23771 1 5

*10 delete*

*DO NOT use a template.*

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/718,321

DATE: 05/21/2002  
TIME: 14:28:55

Input Set : A:\Cura-99.app  
Output Set: N:\CRF3\05212002\I718321.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 7

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/718,321

DATE: 05/21/2002

TIME: 14:28:55

Input Set : A:\Cura-99.app

Output Set: N:\CRF3\05212002\I718321.raw

L:659 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:35  
L:676 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:36  
L:693 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:37  
L:710 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:38  
L:2177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:120 after pos.:0  
L:23687 M:252 E: No. of Seq. differs, <211> LENGTH:Input:12 Found:11 SEQ:1405  
L:23771 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1411